



# United States Patent and Trademark Office

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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO	
09/846,512	05/01/2001	Rachel Meyers	MP100-185P1R2M	5523	
75	90 04/01/2005		EXAM	INER	
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MILLENNIUM 40 LANDSDO	I PHARMACEUTICALS II WNE STREET	ART UNIT	PAPER NUMBER		
CAMBRIDGE,	MA 02139	1642			
			DATE MAILED: 04/01/2005	5	

Please find below and/or attached an Office communication concerning this application or proceeding.

		Applica	tion No.	Applicant(s)								
	Office Action Summary 09/846,512 MEYERS ET AL.											
Offic	ce Action Summary	Examin	er	Art Unit								
		MISOOF	(YU, Ph.D.	1642								
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Status												
1)⊠ Respons	sive to communication(s) filed	on <u>11 January</u> 20	<u>05</u> .									
2a) ☐ This act	☐ This action is <b>FINAL</b> . 2b)⊠ This action is non-final.											
	<u>,                                    </u>											
Disposition of Claims												
4a) Of th 5)⊠ Claim(s) 6)⊠ Claim(s) 7)□ Claim(s)	4) ☐ Claim(s) 83-104 is/are pending in the application. 4a) Of the above claim(s) is/are withdrawn from consideration.  5) ☐ Claim(s) 94 and 96-104 is/are allowed.  6) ☐ Claim(s) 83-93 and 95 is/are rejected.  7) ☐ Claim(s) is/are objected to.  8) ☐ Claim(s) are subject to restriction and/or election requirement.											
Application Pape	rs											
9)∐ The spec	cification is objected to by the	Examiner.										
10)∐ The draw	ving(s) filed on is/are:	a) accepted or t	o)⊡ objected to by the E	xaminer.								
	t may not request that any objecti			• •								
	nent drawing sheet(s) including to or declaration is objected to b											
Priority under 35	U.S.C. § 119											
Priority under 35 U.S.C. § 119  12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).  a) All b) Some * c) None of:  1. Certified copies of the priority documents have been received.  2. Certified copies of the priority documents have been received in Application No  3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).  * See the attached detailed Office action for a list of the certified copies not received.												
Attachment(s)												
1) Notice of Refere 2) Notice of Draftsr	nces Cited (PTO-892) Derson's Patent Drawing Review (PTC	0.048)	4) Interview Summary ( Paper No(s)/Mail Da									
	losure Statement(s) (PTO-1449 or P		5) Notice of Informal Pa		nent)							

#### **DETAILED ACTION**

#### Continued Examination Under 37 CFR 1.114

A request for continued examination under 37 CFR 1.114, including the fee set forth in 37 CFR 1.17(e), was filed in this application after final rejection. Since this application is eligible for continued examination under 37 CFR 1.114, and the fee set forth in 37 CFR 1.17(e) has been timely paid, the finality of the previous Office action has been withdrawn pursuant to 37 CFR 1.114. Applicant's submission filed on 1/11/2005 has been entered. Since all the previously rejected claims are cancelled, the rejection set forth in the prior Office action is moot.

Claim 83-104 are pending and examined on merits.

The text of those sections of Title 35, U.S. Code not included in this action can be found.

### Claim Rejections - 35 USC § 112

Claims 83-93, and 95 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the written description requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

This new matter rejection is made because of the new limitation "a polypeptide which is at least 95 % identical" to SEQ ID NO:2 or "a polypeptide encoded by a nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to the nucleotide sequence of SEQ ID NO:1 or 3", wherein the claimed polypeptide has

protease activity in base claim 83. The specification as originally filed does not reasonably communicate that the claimed invention is a method using a polypeptide which is at least 95 % identical to SEQ ID NO:2 or a polypeptide encoded by a nucleic acid molecule comprising a nucleotide sequence which is at least 95% identical to the nucleotide sequence of SEQ ID NO:1 or 3, wherein the claimed polypeptide has protease activity. This new matter rejection is also made for claim 95, drawn to "a membrane-bound form of an isolated polypeptide".

Applicant states that support for the new claims are found in the previously presented claims, and pages 49-92 of the specification. However, the Office could not find support in the specification originally filed. Applicant is kindly requested to point out the support for the rejected limitation in the specification as originally filed since the support is not apparent to the Office.

## Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(e) the invention was described in a patent granted on an application for patent by another filed in the United States before the invention thereof by the applicant for patent, or on an international application by another who has fulfilled the requirements of paragraphs (1), (2), and (4) of section 371(c) of this title before the invention thereof by the applicant for patent.

Claims 83-87, 89-91, and 93 are rejected under 35 U.S.C. 102(e) as being anticipated by U.S. Pat. 6,294,663 (the '663 patent, issue, filed 02 March 2000).

Claims 83-87, 89-91, and 93 have two part active steps, i.e. contacting SEQ ID NO: 2 protein with a compound (as listed in claims 87, and 89) and determine whether

the compound interacts with the protein, wherein the interaction is determined by the various art-known method listed in claim 93.

The '663 patent discloses a protein sequence (i.e. SEQ ID NO: 2) that is at least 95 % identical to instant SEQ ID NO: 2 (note Exhibit A, the attached sequence alignment). The instant claims read on the immunohistochemstry assay of the prior art at Fig. 7.

# Allowable Subject Matter

Claims 94, and 96-104 are allowed.

#### Conclusion

Any inquiry concerning this communication or earlier communications from the examiner should be directed to MISOOK YU, Ph.D. whose telephone number is 571-272-0839. The examiner can normally be reached on 8 A.M. to 5:30 P.M., every other Friday off.

Any inquiry of a general nature, matching or filed papers or relating to the status of this application or proceeding should be directed to the Judy Ladringan for Art Unit 1642 whose telephone number is 571-272-0536.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Jeffrey Siew can be reached on 571-272-0787. The fax phone number for the organization where this application or proceeding is assigned is 703-872-9306.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR.

Application/Control Number: 09/846,512

Art Unit: 1642

Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

MISOOK YU, Ph.D. Examiner
Art Unit 1642

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APPLICANT: Underwood, Lowell J.
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NUMBER OF SEQ ID NOS: 33
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                                                                                         ; TYPE: PRT
; ORGANISM: Homo
US-09-342-749-2
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APPLICANT: Underwood, Lowell J.
APPLICANT: Underwood, Lowell J.
TITLE OF INVENTION: Transmembrane Serine Protease Overexpressed
TITLE OF INVENTION: in Ovarian Carcinoma and Uses Thereof
FILE REFERENCE: D6192CIP
CURRENT APPLICATION NUMBER: US/09/518,046
CURRENT FILING DATE: 2000-03-02
EARLIER APPLICATION NUMBER: 09/261,416
EARLIER APPLICATION NUMBER: 09/261,416
EARLIER APPLICATION NUMBER: 09/261,416
EARLIER OF SEG ID NOS: 153
NUMBER OF SEG ID NOS: 153
LENGTH: 294
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09342749
Patent No. 6166194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/09518046
Query Match 36.2
Best Local Similarity 46.0
Matches 189, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/342,749
CURRENT FILING DATE: 1999-06-29
                                                                                                                                                                                                                                                                                                                                  APPLICANT: Myriad Genetics, Inc.
TITLE OF INVENTION: TMPRSS2 is a Tumor Suppressor
FILE REPERENCE: 2318-202
                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT: Wong, Alexander K.C.
PPLICANT: Tavtigian, Sean V.
PPLICANT: Teng, DAvid H.-F.
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Local Similarity 88.1%;
tee 258; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     241 HLCGGSVITPLWIITAAHCVYEIVAPRERADRRGRKLLCWRKPTKMKGPRPSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     240 HLCGGSVITPLMIITAAHCVYDLYLPKSWTIQVGLVSL-----LDNPAPSH 285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         180 KVTALHHSVYVREGCASGHVVTIQCTACGHRRGYSSRIVGGNMSLLSQWPWQASLQPQGY 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121 PTAASWKTMCSDDWKGHYANVACAQLGPPSYVSSDNLRVSSLEGQPREEFVSIDHLLPDD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FTAASWKTMCSDDWKGHYANVACAQLGPPSYVSSDNLRVSSLEGQFREEFVSIDHLLPDD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MGENDPPAVEAPFSFRSLFGLDDLKISFVAPDADAVAAQILSLLPLKFF-PIIVIGIIAL
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                                                                                                          sapiens
  36.2%; Score 885; DB 3; Length 492.
46.0%; Pred. No. 1.4e-77;
live 53; Mismatches 149; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 1354; Di
Pred. No. 1.6e
7; Mismatches
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20;

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